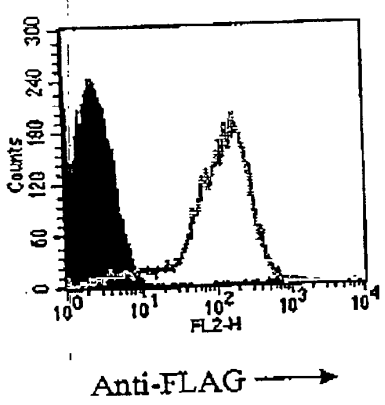

Figure 1

10/070982


```
          10        30        50        70        90
    ACCACTGCTTCATTGCTGCTGAGAATGTTCCAGGCTGAGTGAGAAGTAAAAATTCATCATCTCTGAAGAACTCTTACCCAGCCCTGTTGA
             110       130       150       170

                              M  G  S  R  C  L  L  H  A  S  V  P  G  M  L  L  I  W    18
1   AGAAATTCCCAGAATGTTGATGGGAGCAACTAGAGATATGGGAAGCAGGTGCCTTCTGCATGCCTCAGTTCCTGGAATGCTGCTTATCTG
       190       210       230       250       270

19  S  I  L  Q  M  M  N  I  S  A  S  C  P  Q  C  N  E  N  A  S  C  F  N  S  T  H  C  V  C  K    48
    GTCAATATTACAAATGATGAATATTTCAGCTTCCTGTCCCCAGTGCAATGAAAATGCCAGCTGCTTCAACAGCACCCACTGTGTTTGTAA
          290       310       330       350

49  E  G  F  W  T  G  S  E  N  R  R  I  I  E  P  H  E  K  C  Q  D  I  N  E  C  L  L  K  E  L    78
    AGAAGGATTCTGGACGGGCTCTGAGAATAGAAGAATTATTGAGCCCCATGAGAAATGTCAAGATATTAATGAGTGTCTACTGAAAGAATT
       370       390       410       430       450

79  V  C  K  D  V  S  Y  C  R  N  K  I  G  T  Y  I  C  S  C  V  V  K  Y  P  L  F  N  W  V  A    108
    GGTATGCAAGGATGTGTCGTACTGCAGAAATAAAATTGGGACTTACATATGCAGCTGTGTAGTAAAATATCCTTTGTTCAACTGGGTAGC
          470       490       510       530

109 G  I  I  N  I  D  H  P  D  C  Y  V  N  K  S  K  N  T  G  S  K  T  H  T  L  G  V  L  S  E    138
    TGGCATTATTAATATTGATCACCCTGATTGTTATGTGAACAAGAGCAAGAATACAGGATCAAAAACACATACTTTGGGAGTACTGAGTGA
       550       570       590       610       630

139 F  K  S  K  E  E  V  A  K  G  A  T  K  L  L  R  K  V  E  H  H  I  L  N  E  N  S  D  I  P    168
    ATTTAAATCCAAAGAGGAGGTTGCAAAAGGAGCTACCAAGTTACTTCGCAAAGTGGAACATCACATCTTGAATGAAAACTCAGATATACC
          650       670       690       710

169 K  K  D  E  N  P  L  L  D  I  V  Y  E  T  K  R  C  K  T  M  T  L  L  E  A  G  N  N  T  M    198
    AAAAAAGGATGAAAATCCTTTATTGGATATAGTGTATGAAACTAAGAGGTGCAAGACGATGACTCTTCTAGAAGCTGGCAACAACACAAT
       730       750       770       790       810

199 K  V  D  C  T  S  G  F  K  E  H  N  S  G  G  E  T  A  V  A  F  I  A  Y  K  S  L  G  N  L    228
    GAAGGTTGACTGCACTAGTGGTTTCAAAGAGCACAACAGTGGAGGTGAAACTGCAGTGGCTTTCATTGCATATAAGTCTCTTGGGAATCT
          830       850       870       890

229 L  N  G  S  F  F  S  N  E  E  G  F  Q  E  V  T  L  N  S  H  I  V  S  G  A  I  R  S  E  V    258
    TCTAAATGGTTCCTTTTTTAGTAATGAAGAAGGGTTTCAGGAAGTGACACTGAACTCTCACATCGTTAGTGGAGCCATTCGCTCAGAGGT
```

Figure 2

10/070982


```
      910                 930                 950                 970                 990
259   K  P  V  L  S  E  P  V  L  L  T  L  Q  N  I  Q  P  I  D  S  R  A  E  H  L  C  V  H  W  E   288
      CAAACCTGTCCTCTCTGAACCTGTACTCCTGACTTTACAAAATATTCAGCCCATTGACTCAAGAGCAGAACATCTCTGTGTCCATTGGGA
                1010                1030                1050                1070

289   G  S  E  E  G  G  S  W  S  T  K  G  C  S  H  V  Y  T  N  N  S  Y  T  I  C  K  C  F  H  L   318
      AGGATCAGAGGAAGGGGGGAGCTGGTCTACCAAAGGATGCTCTCACGTGTACACCAATAATTCCTACACCATTTGCAAGTGTTTCCACCT
          1090                1110                1130                1150                1170

319   S  S  F  A  V  L  M  A  L  P  H  E  E  D  G  V  L  S  A  L  S  V  I  T  Y  V  G  L  S  L   348
      GTCCAGCTTTGCTGTGCTCATGGCTCTACCCCATGAGGAGGATGGTGTGCTTTCTGCACTCTCTGTGATCACCTATGTGGGACTGAGTCT
                1190                1210                1230                1250

349   S  L  L  C  L  F  L  A  A  I  T  F  L  L  C  R  P  I  Q  N  T  S  T  T  L  H  L  Q  L  S   378
      TTCTCTCTTGTGCCTATTTCTGGCGGCCATCACTTTTCTCCTGTGCCGACCCATTCAGAATACCAGCACGACACTCCACCTGCAGCTCTC
          1270                1290                1310                1330                1350

379   I  C  L  F  L  A  D  L  L  F  L  T  G  I  N  R  T  K  P  K  V  L  C  S  I  I  A  G  M  L   408
      CATCTGCCTTTTCCTGGCTGACCTCCTCTTCCTCACAGGCATCAACAGAACTAAGCCTAAGGTGCTGTGCTCCATCATAGCGGGGATGTT
                1370                1390                1410                1430

409   H  Y  L  Y  L  A  S  F  M  W  M  F  L  E  G  L  H  L  F  L  T  V  S  N  L  K  V  A  N  Y   438
      GCACTACCTCTACTTGGCTTCCTTCATGTGGATGTTTCTGGAAGGGCTACATCTTTTTCTCACTGTGAGCAATCTCAAAGTGGCCAACTA
          1450                1470                1490                1510                1530

439   S  N  S  G  R  F  K  K  R  F  M  Y  P  V  G  Y  G  L  P  A  F  I  V  A  V  S  A  I  A  G   468
      CAGCAACTCAGGCAGATTCAAGAAGAGGTTCATGTATCCTGTAGGATATGGGCTTCCTGCTTTTATTGTTGCTGTATCTGCAATAGCTGG
                1550                1570                1590                1610

469   H  K  N  Y  G  T  H  N  H  C  W  L  S  L  H  R  G  F  I  W  S  F  L  G  P  A  A  A  I  I   498
      CCACAAGAATTATGGAACACACAACCACTGCTGGCTCAGCCTTCATCGAGGATTCATCTGGAGCTTCTTGGGGCCAGCGGCAGCCATTAT
          1630                1650                1670                1690                1710

499   L  I  N  L  V  F  Y  F  L  I  I  W  I  L  R  S  K  L  S  S  L  N  K  E  V  S  T  L  Q  D   528
      CTTGATAAACCTGGTGTTCTACTTTCTAATAATATGGATTTTGAGAAGCAAACTTTCTTCTCTCAATAAAGAAGTTTCTACACTTCAAGA
                1730                1750                1770                1790

529   T  K  V  M  T  F  K  A  I  V  Q  L  F  V  L  G  C  S  W  G  I  G  L  F  I  F  I  E  V  G   558
      CACAAAGGTTATGACATTTAAAGCCATTGTCCAGTTATTTGTGTTGGGATGTTCTTGGGGCATTGGCTTGTTTATTTTCATTGAAGTTGG
```

Figure 2 (continued)

10/070982

```
     1810          1830          1850          1870          1890
559  K  T  V  R  L  I  V  A  Y  L  F  T  I  I  N  V  L  Q  G  V  L  I  F  M  V  H  C  L  L  N   588
GAAGACAGTGAGACTGATCGTTGCCTATCTGTTCACCATCATCAATGTCCTGCAGGGTGTTTTGATATTTATGGTACATTGTCTGCTTAA
          1910          1930          1950          1970

589  R  Q  V  R  M  E  Y  K  K  W  F  H  R  L  R  K  E  V  E  S  E  S  T  E  V  S  H  S  T  T   618
TCGCCAGGTGCGGATGGAATATAAGAAGTGGTTTCATAGACTGCGGAAGGAAGTTGAAAGTGAAAGCACTGAAGTGTCTCATTCTACTAC
    1990          2010          2030          2050          2070

619  H  T  K  M  G  L  S  L  N  L  E  N  F  C  P  T  G  N  L  H  D  P  S  D  S  I  L  P  S  T   648
TCACACAAAAATGGGTCTTTCTCTGAACCTGGAAAATTTCTGCCCAACAGGAAACCTCCATGATCCTTCTGACTCCATCCTTCCAAGTAC
          2090          2110          2130          2150

649  E  V  A  G  V  Y  L  S  T  P  R  S  H  M  G  A  E  D  V  N  S  G  T  H  A  Y  W  S  R  T   678
TGAAGTAGCAGGTGTATATCTAAGCACACCCAGGTCTCACATGGGTGCTGAGGATGTGAACTCAGGTACTCACGCTTACTGGAGCAGAAC
    2170          2190          2210          2230          2250

679  I  S  D  *                                                                                681
TATTAGTGATTGAATCAGCTCCTTCCCCCAAGCCTCTTACAGTACATTTTAACTTGTACTGTGCCATGCACATGAAGCTATAATTGCTAG
          2270          2290          2310          2330

TCTGGTAAAACAACTGTTGCATATTCCATGATCATTTCATTTTATCTCTACTTGCAAAAGTTAGCTTTCTTTTTATATCATTTTTATTTC
    2350          2370          2390          2410          2430

TCTTTCTTTTGTTTATATATAGCTTCAGTTGAGTGGGTTTCTAGTCTTAATGTTCTAGATCACTATTTTCTTTTCAGTTAACCTTTATTG
          2450          2470          2490          2510

GTATTTAGTTCCTGTGTAGTGTATACCACTGGAATATTTTTATTTCTTTAATTTTGAGGTTAAAATATAGTTACATCATTTTTCCTTTTT
    2530          2550          2570          2590          2610

TTCTTTCCCACAATCCTCCTGTATACTTTTTCCCTGGTGTCTATTTTATTGTTTCTACATGCATATATATTTTATGCAAAACATATATAT
          2630          2650          2670          2690

GTATAAATATAAATATATATTCTTATATGCATGAAAACCATCTACTTCATCCAAATAATGTTCCTTCTATGTATGTTTTCAGGACAGGGA
    2710          2730          2750          2770          2790

CAACAATAGCTATGGTAGCATGGCAGGGGAAAGCCCACAGGACCTCAGCCTTATACAAAGAATCAGAGGCAACTGAGGAGTGCTGAGTTG
          2810          2830          2850          2870

AAGGAATTGTCTTACCCAGGGGAGGGCACATTAATTGGTTATCTAATACAAAATGTTCAGCCCCAAAACTGTTAAGATAAAAGCCTATAT
```

Figure 2 (continued)

```
2890                2910                2930                2950                2970
GCATCTTAGGAAGTATCTACCTTGATACACCTTTATTGGAATATCATCCACATGTTTATTGTGTGTTCTGAAGAGGGTCTGTTGAATTTC
                    2990                3010                3030                3050
TAAGGGTTGATCAGTTTAATTGTGCCATTTTATATTCAGGGTGTTTGGCTTTGTTGTAGTGAATAATGCTATATTTCCCTGTATGTGTCA
      3070                3090                3110                3130                3150
TCTTTGACTGTTATTTTTTCCTGGCGATACTTTATTCAACAAGAACCTAGAGCCTTGGTTTATTACTTTTTCTTCCATAGAAAAACTATT
                3170                3190                3210                3230
TGTCTTCCAGGATTAGATATGATCAATATTTCTTATATGCATGTATCAAATATCATGATGAAATATATTACTGTGTATAATTAATAACTG
      3250
GCAATAAAGTCGAAGGGA
```

Figure 2 (continued)

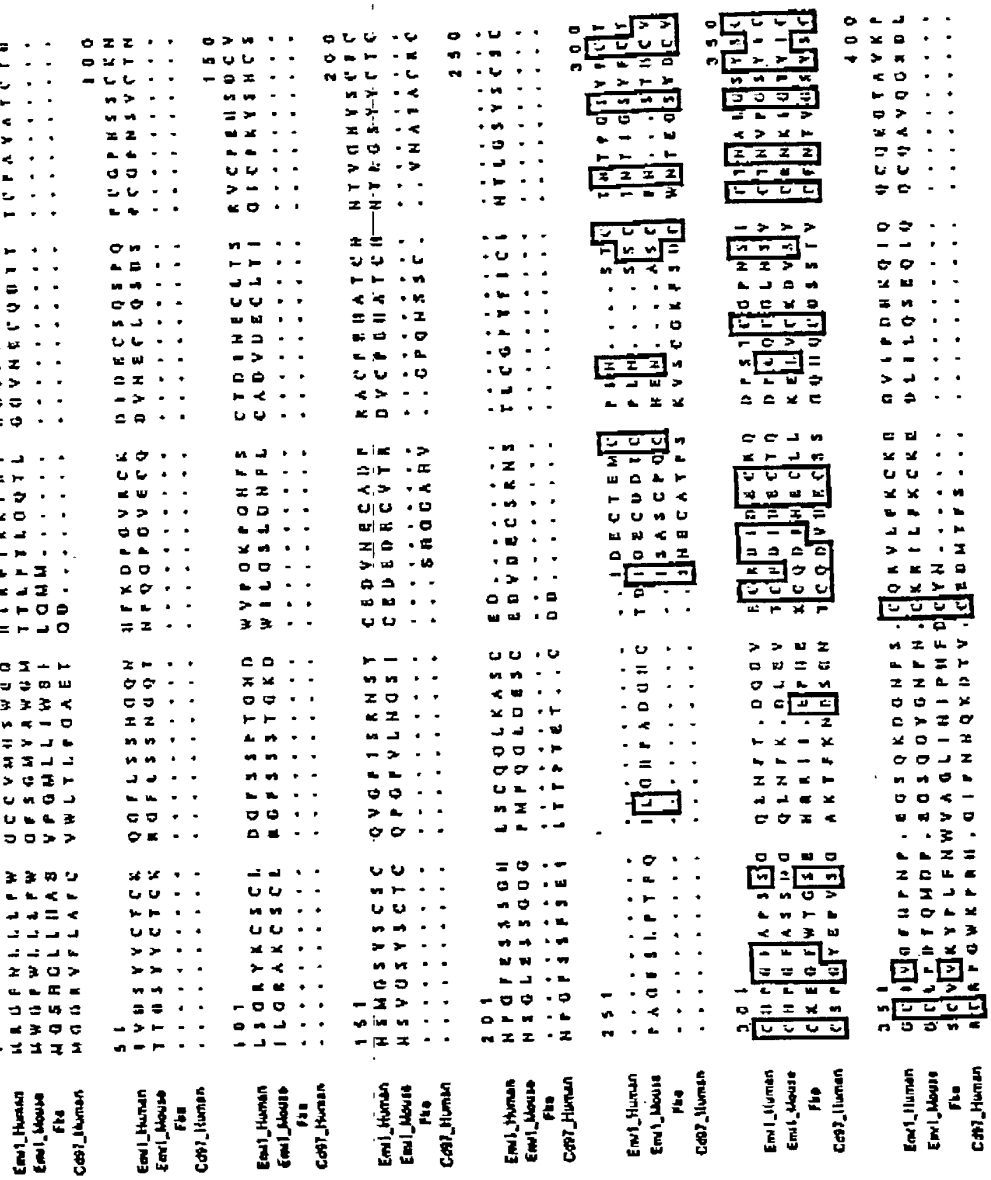

Figure 3

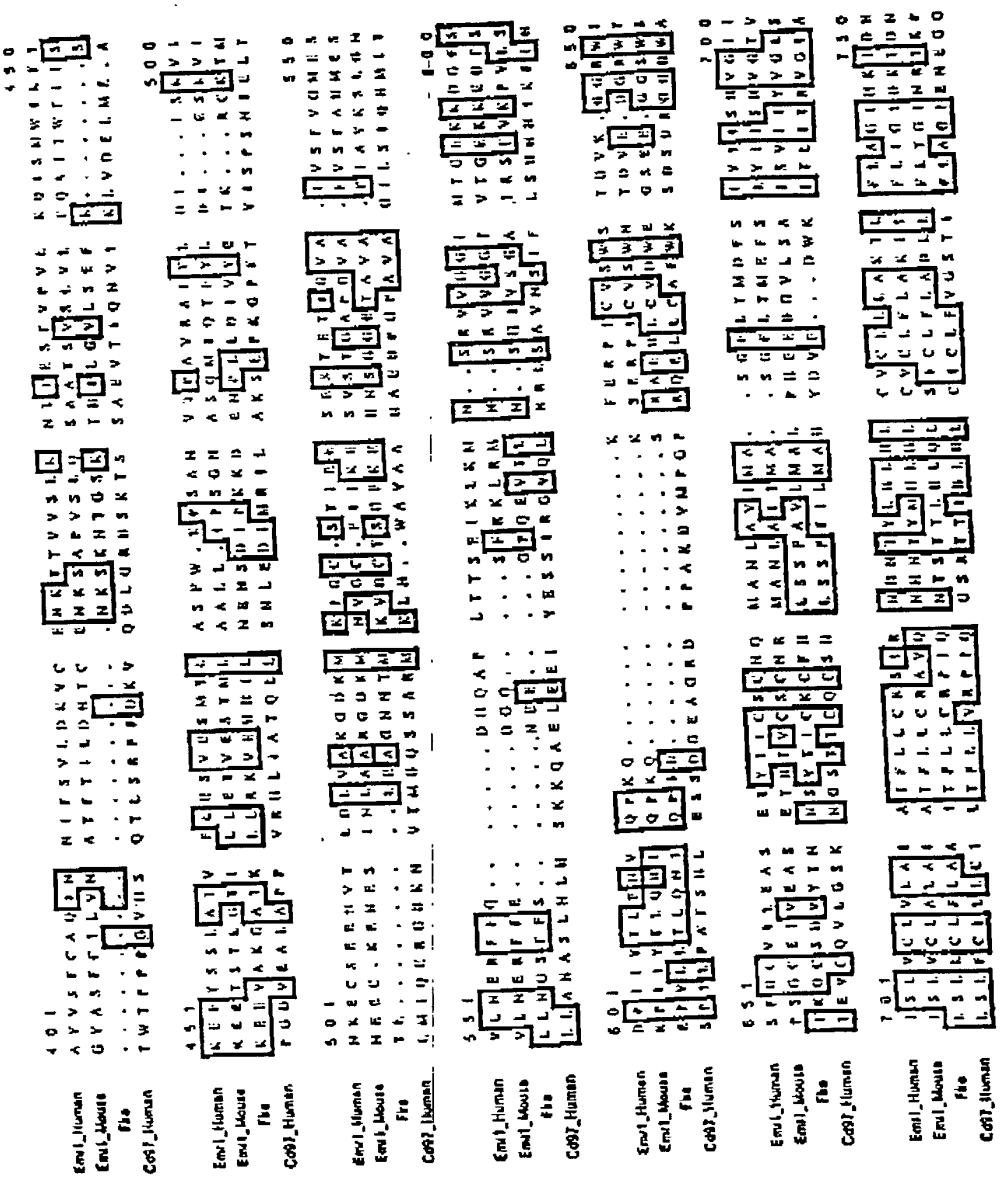

Figure 3 (continued)

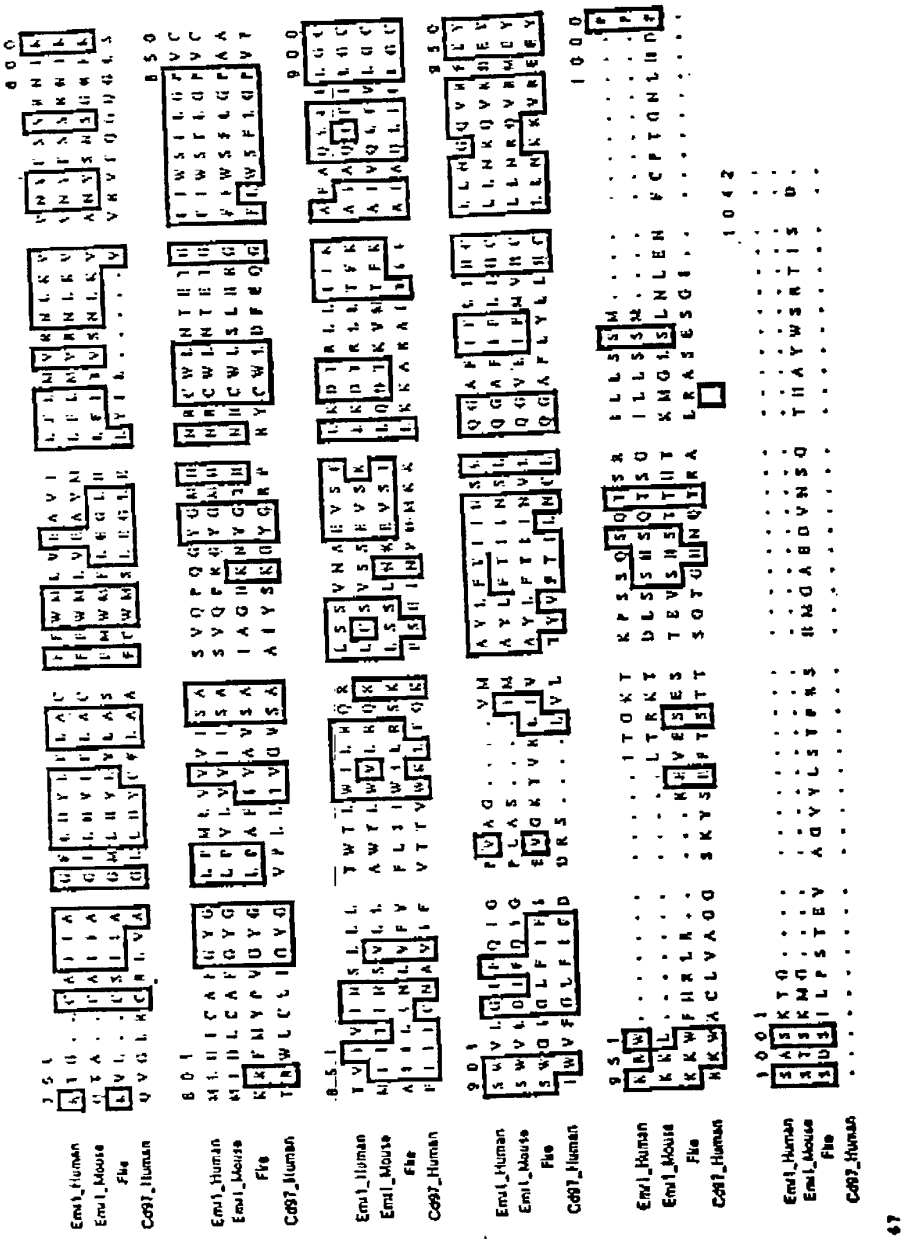

Figure 3 (continued)

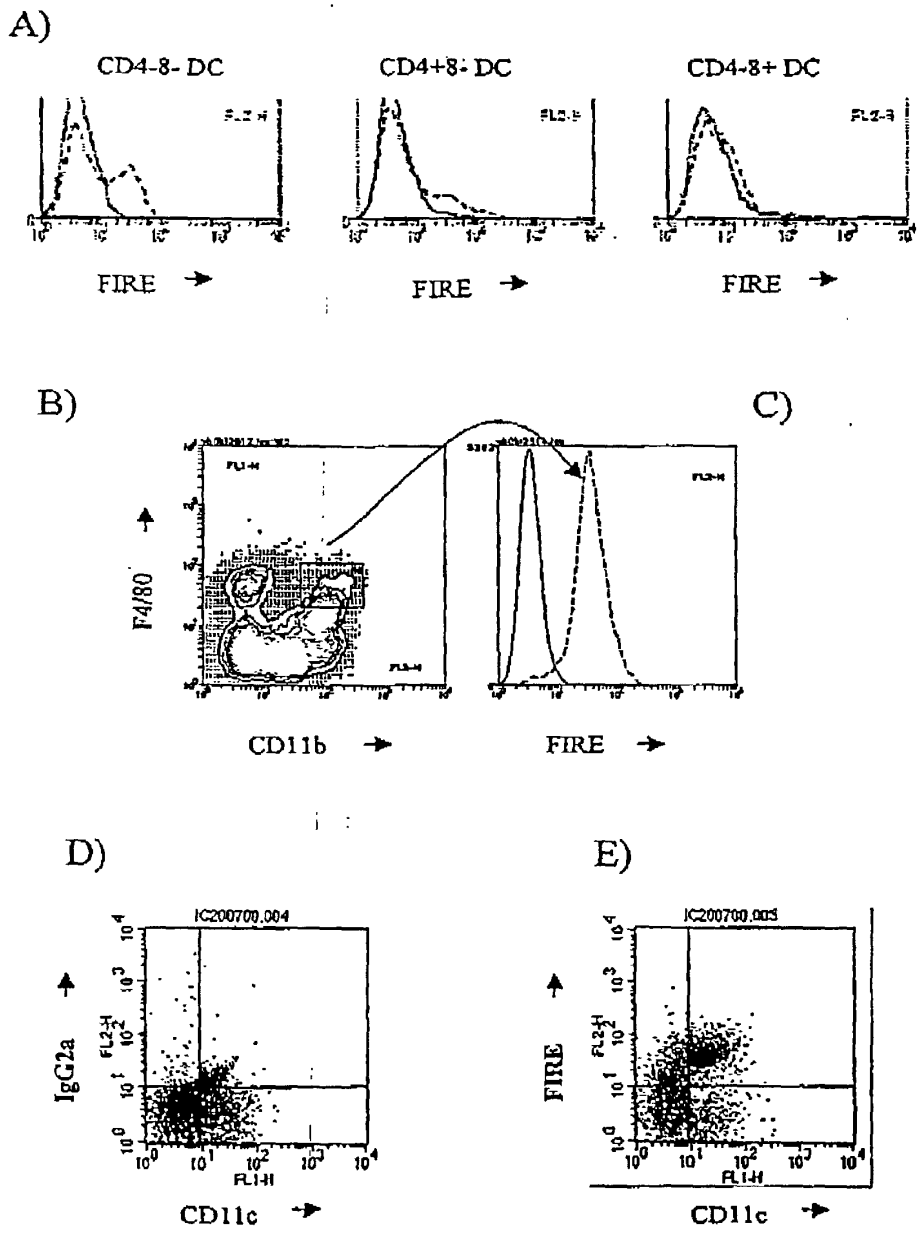
A)
CD4-8- DC
CD4+8- DC
CD4-8+ DC
FIRE
FIRE
FIRE
B)
C)
F4/80
CD11b
FIRE
D)
E)
IC200700.004
IC200700.005
IgG2a
FIRE
CD11c
CD11c

Figure 4

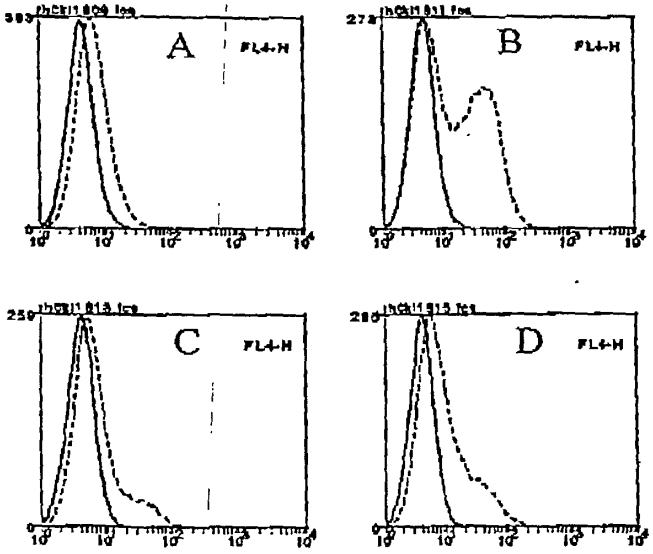

Figure 5

```
        A  S  C  P  P  C  P  K  Y  A  S  C  H  N  S  T  H  C  T  C  E  D  G  F  R  A  R  S  G  R
   1    GCTTCCTGTCCTCCATGCCCTAAATATGCCAGCTGCCACAACAGCACCCACTGTACTTGTGAAGATGGCTTTCGGGCCAGGTCTGGCAGG    90
        T  Y  F  H  D  S  S  E  K  C  E  D  I  N  E  C  E  T  G  L  A  K  C  K  Y  K  A  Y  C  R
  91    ACATACTTTCATGATTCCTCTGAGAAGTGTGAAGATATTAATGAATGTGAAACCGGGCTGGCAAAGTGCAAGTATAAAGCATATTGTAGG    180
        N  K  V  G  G  Y  I  C  S  C  L  V  K  Y  T  L  F  N  F  L  A  G  I  I  D  Y  D  H  P  D
 181    AATAAAGTTGGAGGTTACATCTGTAGCTGTTTGGTAAAATATACTTTATTCAACTTTCTGGCTGGTATTATAGATTATGATCATCCGGAT    270
        C  Y  E  N  N  S  Q  G  T  T  Q  S  N  V  D  I  W  E  N  L  R  R  N  G  S  R  E  D  F  A
 271    TGTTACGAGAACAATAGTCAAGGGACGACACAGTCAAACGTGGATATTTGGGAAAATCTGAGAAGAAATGGAAGCAGAGAGGACTTTGCA    360
        R  R  A  T  Q  L  I  Q  S  V  E  L  S  I  W  N  A  S  F  A  S  P  G  K  G  Q  I  S  E  F
 361    AGAAGGGCTACTCAACTAATTCAAAGCGTGGAGTTGAGCATCTGGAATGCGAGTTTTGCTTCTCCAGGAAAGGGTCAAATTTCTGAATTT    450
        D  I  V  Y  E  T  K  R  C  N  E  T  R  E  N  A  F  L  E  A  G  N  N  T  M  D  I  N  C  A
 451    GATATAGTCTATGAAACCAAGAGGTGCAATGAGACAAGGGAGAATGCTTTTCTGGAAGCTGGAAATAACACCATGGATATCAACTGTGCT    540
        D  A  L  K  G  N  L  R  E  S  T  A  V  A  L  I  T  Y  Q  S  L  G  D  I  L  N  A  S  F  F
 541    GATGCTTTAAAAGGAAACCTAAGAGAGAGCACTGCAGTTGCCCTAATCACTTATCAATCTCTTGGGGATATTCTGAATGCATCCTTTTTT    630
        S  K  R  K  G  M  Q  E  V  K  L  N  S  Y  V  V  S  G  T  V  G  L  K  E  K  I  S  L  S  E
 631    AGTAAACGAAAAGGGATGCAGGAAGTAAAACTGAACTCTTACGTTGTGAGCGGCACCGTCGGTTTGAAGGAAAAAATTTCCCTCTCTGAA    720
        P  V  F  L  T  F  R  H  N  Q  P  G  D  K  R  T  K  H  I  C  V  Y  W  E  G  S  E  G  G  R
 721    CCTGTGTTCCTGACTTTTCGCCATAATCAGCCTGGTGACAAGAGAACAAAACATATCTGTGTCTACTGGGAGGGATCAGAGGGAGGCCGC    810
        W  S  T  E  G  C  S  H  V  H  S  N  G  S  Y  T  K  C  K  C  F  H  L  S  S  F  A  V  L  V
 811    TGGTCCACGGAGGGCTGCTCTCATGTGCACAGCAACGGTTCTTACACCAAATGCAAGTGCTTCCATCTGTCCAGCTTTGCCGTCCTCGTG    900
        A  L  A  P  K  E  D  P  V  L  T  V  I  T  Q  V  G  L  T  I  S  L  L  C  L  F  L  A  I  L
 901    GCTCTTGCCCCCAAGGAGGACCCTGTGCTGACCGTGATCACCCAGGTGGGGCTGACCATCTCYCTGCTGTGCCTCTTCCTGGCCATCCTC    990
        T  F  L  L  C  R  P  I  Q  N  T  S  T  S  L  H  L  E  L  S  L  C  L  F  L  A  H  L  L  F
 991    ACCTTCCTCCTGTGCCGGCCCATCCAGAACACCAGCACCTCCCTCCATCTAGAGCTCTCCCTCTGCCTCTTCCTGGCCCACCTCCTGTTC    1080
        L  T  G  I  N  R  T  E  P  E  V  L  C  S  I  I  A  G  L  L  H  F  L  Y  L  A  C  F  T  W
1081    CTGACGGGCATCAACAGAACTGAGCCTGAGGTGCTGTGCTCCATCATTGCAGGGCTGCTGCACTTCCTCTACCTGGCTTGCTTCACCTGG    1170
        M  L  L  E  G  L  H  L  F  L  T  V  R  N  L  K  V  A  N  Y  T  S  T  G  R  F  K  K  R  F
1171    ATGCTCCTGGAAGGGCTGCACCTCTTCCTCACCGTCAGGAACCTCAAGGTGGCCAACTACACCAGCACGGGCAGATTCAAGAAGAGGTTC    1260
        M  Y  P  V  G  Y  G  I  P  A  V  I  I  A  V  S  A  I  V  G  P  Q  N  Y  G  T  F  T  H  C
1261    ATGTACCCTGTAGGCTACGGGATCCCAGCTGTGATTATTGCTGTGTCAGCAATAGTTGGACCCCAGAATTATGGAACATTTACTCACTGT    1350
        W  L  K  L  D  K  G  F  I  W  S  F  M  G  P  V  A  V  I  I  L  I  N  L  V  F  Y  F  Q  V
1351    TGGCTCAAGCTTGATAAAGGATTCATCTGGAGCTTCATGGGGCCAGTAGCAGTCATTATCTTGATAAACCTGGTGTTCTACTTCCAAGTT    1440
        L  W  I  L  R  S  K  L  S  S  L  N  K  E  V  S  T  I  Q  D  T  R  V  M  T  F  K  A  I  S
1441    CTGTGGATTTTGAGAAGCAAACTTTCCTCCCTCAATAAAGAAGTTTCCACCATTCAGGACACCAGAGTCATGACATTTAAAGCCATTTCT    1530
        Q  L  F  I  L  G  C  S  W  G  L  G  F  F  M  V  E  E  V  G  K  T  I  G  S  I  I  A  Y  S
1531    CAGCTATTTATCCTGGGCTGTTCTTGGGGCCTTGGTTTTTTATGGTTGAAGAAGTAGGGAAGACGATTGGATCAATCATTGCATACTCA    1620
```

Figure 6

10/070982

```
      F  T  I  I  N  T  L  Q  G  V  L  L  F  V  V  H  C  L  L  N  R  Q  V  R  M  E  Y  K  K  W
1621  TTCACCATCATCAACACCCTTCAGGGAGTGTTGCTCTTTGTGGTACACTGTCTCCTTAATCGCCAGGTTCGAATGGAATATAAAAAGTGG  1710
      F  S  G  M  R  K  G  V  E  T  E  S  T  E  M  S  R  S  T  T  Q  T  K  T  E  E  V  G  K  S
1711  TTTAGTGGGATGCGGAAAGGGGTAGAAACTGAAAGCACTGAGATGTCTCGCTCTACTACCCAAACCAAAACGGAAGAAGTGGGGAAGTCC  1800
      S  E  I  F  H  K  G  G  T  A  S  S  S  A  E  S  T  K  Q  P  Q  P  Q  V  H  L  V  S  A  A
1801  TCAGAAATCTTTCATAAAGGAGGCACTGCATCATCATCTGCAGAGTCAACCAAGCAACCGCAGCCACAGGTTCATCTCGTCTCTGCTGCT  1890
      W  L  K  M  N  *
1891  TGGCTAAAGATGAACTGACCTGGCAAGTGCCATGGCAATGACCCGGAAGTTACCGCTCCTTTCCGTTTGTCTACAGCGCCCCTGTGGTCA  1980
1981  CACATAGATTGGACAAATGCCACTATTTCTAGCTTTCCTGTGAAAAGTCTAGGCTCATTCACCTATTTTGGCTTTTTATGTTCATAGAAA  2070
2071  GAACAAGACATTTGGGAGAATTCTTAGATCCAGAGTCCAGTAGTGTGGCACGTGCAATGAAGTGTCGGAAGGATGCATTTTAAAGATGGC  2160
2161  GGGCGGGAGAAGTGGATTTTTCTTCTTGCAGCTACTGCCACCTTGCCAGAAACTTCACTAACTGGCATCTGGRATTCAGCTCATAGTTCC  2250
2251  CTTTCTGGCCTCTCTGCTGTATTTTATGCTCCCAAAGATCTTACATTAACACTCCACATTCACATAATTCAACAATTTTCATATGGATCA  2340
2341  GTATTAAAGAGGGTGTTGCATTTTGCAATACAAAAATGCATTATCAGGTGCTGGAGAGGATGTGGAGAAATAGGAACACTTTTACACTGT  2430
2431  TGGTGGGACTGTAAACTAGTTCAACCATCGTGGAAGTCAGTGTGGCGATTCCTCAGGGATCTAGAACTAGAAATACCATTTGACACAGCT  2520
2521  ATCCCATTACTGGGTATATACCCAAAGGACTATAAATCATGCTGCTATAAAGACACATGCACACGTATGTTTATTGTGGCATTATTCACA  2610
2611  ATAGCAAAGACTTGGAACCAACCCAAATGTCCAACAATGATAGACTGGATTAAGAAAATGTGGCACATATACACCATGGAATACTATGCA  2700
2701  GCCATAAAAAATGATGAGTTCATGTCCTTTGTAGGGACATGGATGAAATCCGGAATTCCGCCGATACTGACGGGCTCCAGGAGTCGTCGC  2790
2791  CACCAATC  2798
```

Figure 6 (continued)